## STIC REPORT 10/017,870 034827-1401

```
SEQ ID NO: 1
RESULT 3
US-08-938-669A-2
; Sequence 2, Application US/08938669A
; Patent No. 5171788
  GENERAL INFOFMATION:
     APPLICANT: Nguyen, Thai D.
     APPLICANT: Polansky, Jon R.
     TITLE OF INVENTION: METHODS FOR THE DIAGNOSIS,
TITLE OF INVENTION: PROGNOSIS AND TREATMENT OF GLAUCOMA AND
     TITLE OF INVENTION: RELATED DISEASES
    NUMBER OF SEQUENCES: 32
    CORRESPONDENCE ADDRESS:
      ADDRESSEE: Howrey & Simon
       STREET: 1299 Pennsylvania Avenue, N.W.
;
       CITY: Washington STATE: DC
;
;
       COUNTRY: USA
;
;
       ZIP: 20004-2402
    COMPUTER READABLE FORM:
;
      MEDIUM TYPE: Diskette
;
       COMPUTER: IBM Compatible
;
      OPERATING SYSTEM: DOS
;
       SOFTWARE: FastSEQ for Windows Version 2.0
;
    CUPRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/938,669A
       FILING DATE:
;
      CLASSIFICATION: 435
;
    PRIOR APPLICATION DATA:
      APPLICATION NUMBER: 08/791,154
       FILING DATE: 28-JAN-1997
;
    ATTOPHEY/AGENT INFOFMATION:
      NAME: Mendelson, Elliot
       PEGISTRATION NUMBER: F-42,878
      FEFERENCE/DOCKET NUMBER: 07425~0034
    TELECOMMUNICATION INFOFMATION:
       TELEPHONE: 202 383+€857
       TELEFAM: 202 383-6610
       TELEM:
 INFORMATION FOR SEQ ID NO:
   SEQUENCE CHAPACTERISTICS:
       LENGTH: 5304 base pairs
       TYPE: nucleic acid
       STFANDEDNESS: single
       TOPOLOGY: linear
US-08-938-669A-2
  Query Match
                           100.0-; Score 32; DB 4; Length 5304;
  Best Local Similarity 100.0-; Pred. No. 1.3e-09; Matches 32; Conservative 0; Mismatches 0; Indels
Gaps 0;
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## SEQ ID NO:2

```
RESULT 1
US-08-791-347-9
; Sequence 9, Application US/08791347
; Patent No. 5885776
  GENEFAL INFORMATION:
     APPLICANT: Stone, Edwin M.
     APPLICANT: Sheffield, Val C. APPLICANT: Alward, Wallace L.M.
     TITLE OF INVENTION: GLAUCOMA COMPOSITIONS AND THERAPEUTIC TITLE OF INVENTION: AND DIAGNOSTIC USES THEREFOR
    NUMBER OF SEQUENCES: 16
    COPRESPONDENCE ADDRESS:
      ADDRESSEE: FOLEY, HOAG & ELIOT LLP
      STREET: One Post Office Square
      CITY: Boston
;
      STATE: MA
      COUNTEY: USA
       ZIP: 02109-2170
    COMPUTER READABLE FORM:
     MEDIUM TYPE: Florry disk
      COMPUTER: IBM FC compatible
      OPERATING SYSTEM: PC-DOS/MS-DOS
      SOFTWARE: FatentIn Release #1.0, Version #1.30
    CUFFENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/791,347
      FILING DATE: 30-JAN-1997
      CLASSIFICATION: 514
    ATTORNEY/AGENT INFOFMATION:
      NAME: Arnold, Beth E.
      FEGISTPATION NUMBER: 35,430
      PEFERENCE/DOCKET NUMBER: UIA-010.26
     TELECOMMUNICATION INFORMATION:
       TELEPHONE: 617-832-1000
       TELEFAX: 617-832-7000
   INFORMATION FOR SEQ ID NO:
    SEQUENCE CHARACTERISTICS:
       LENGTH: 190 base pairs
       TYPE: nucleic acid
       STRANDEDNESS: single
       TOPOLOGY: linear
    MOLECULE TYPE: DNA
US-08-791-347-9
                          100.0:; Score 22; DB 2; Length 190;
  Query Match
  Best Local Similarity 100.0-; Pred. No. 0.00047;
 Matches 22; Conservative 0; Mismatches 0; Indels
       0;
Gaps
Qу
        1 CCGTATTCTTGGGGTGGCTACA 22
```

11111111111111111111

Db 160 CCGTATTCTTGGGGTGGCTACA 181

FIGURE B

SEQ ID NO: 3

**RESULT 1** 

US-08-791-347-7

; Sequence 7, Application US/08791347

; Patent No. 5885776

: GENERAL INFORMATION:

; APPLICANT: Stone, Edwin M.

; APPLICANT: Sheffield, Val C.

; APPLICANT: Alward, Wallace L.M.

; TITLE OF INVENTION: GLAUCOMA COMPOSITIONS AND THERAPEUTIC

; TITLE OF INVENTION: AND DIAGNOSTIC USES THEREFOR

; NUMBER OF SEQUENCES: 16

; CORRESPONDENCE ADDRESS:

: ADDRESSEE: FOLEY, HOAG & ELIOT LLP

STREET: One Post Office Square

CITY: Boston

STATE: MA

COUNTRY: USA

ZIP: 02109-2170

; COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

; COMPUTER: IBM PC compatible

; OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

; CURRENT APPLICATION DATA

APPLICATION NUMBER: US/08/791,347

FILING DATE: 30-JAN-1997

CLASSIFICATION: 514

; ATTORNEY/AGENT INFORMATION:

; NAME: Arnold, Beth E.

; REGISTRATION NUMBER: 35,430

; REFERENCE/DOCKET NUMBER: UIA-010.26

: TELECOMMUNICATION INFORMATION:

TELEPHONE: 617-832-1000

; TELEFAX: 617-832-7000

```
: INFORMATION FOR SEQ ID NO: 7:
  SEQUENCE CHARACTERISTICS:
   LENGTH: 195 base pairs
   TYPE: nucleic acid
   STRANDEDNESS: single
   TOPOLOGY: linear
: MOLECULE TYPE: DNA
US-08-791-347-7
 Query Match
                92.3%; Score 24; DB 2; Length 195;
 Best Local Similarity 100.0%; Pred. No. 2.8e-05;
 Matches 24; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
    3 CAAACCTGGGAGACAAACATCCGT 26
Qy
    10 CAAACCTGGGAGACAAACATCCGT 33
Db
FIGURE C
SEQ ID NO 4
RESULT 14
US-08-546-568A-3
; Sequence 3, Application US/08546568A
; GENERAL INFORMATION:
    APPLICANT: NGUYEN, THAI D.
    APPLICANT: POLANSKY, JON R.
    APPLICANT: HUANG, WEIDONG
    TITLE OF INVENTION: METHOD'S FOR THE DIAGNOSIS OF GLAUCOMA
    NUMBER OF SEQUENCES: 3
    COFFESPONDENCE ADDRESS:
     ADDRESSEE: HOWREY & SIMON
      STREET: 1299 PENNSYLVANIA AVE., N.W.
      CITY: WASHINGTON
      STATE: D.C.
      COUNTRY: US
      ZIP: 20004
    COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: IBM PC compatible
      OPERATING SYSTEM: PC-DDS/MS-DOS
      SOFTWARE: PatentIn Release #1.0, Version #1.25
    CURPENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/546,568A
      FILING DATE:
      CLASSIFICATION: 536
   ATTORNEY/AGENT INFORMATION:
    NAME: AUERBACH, JEFFREY I
      REGISTRATION NUMBER: 32,680
    TELECOMMUNICATION INFORMATION:
```

```
TELEPHONE: (202) 383-7451
;
       TELEFAX: (202) 383-6610
;
   INFORMATION FOR SEQ ID NO: 3:
     SEQUENCE CHARACTERISTICS:
;
       LENGTH: 1491 base pairs
;
       TYPE: nucleic acid
;
      STPANDEDNESS: double
      TOFOLOGY: linear
     MOLECULE TYPE: cDNA
     HYPOTHETICAL: NO
     ANTI-SENSE: NO
     IMMEDIATE SOURCE:
       CLONE: TIGH coding sequence
US-08-546-568A-3
  Query Match
                         88.6*; Score 31; DB 9; Length 1491;
  Best Local Similarity 100.0%; Pred. No. 8.1e-07;
 Matches 31; Ccnservative 0; Mismatches 0; Indels
                                                                0;
Gars
       0;
        5 GCTATAAGTACAGCAGCATGATTGACTACAA 35
QУ
          1388 GCTATAAGTACAGCAGCATGATTGACTACAA 1418
FIGURE D
SEQ ID NO: 5
RESULT 3
BM689172
LOCUS
            BM689172
                                    357 bp
                                              mFNA
                                                      linear
                                                               EST 28-
FEB-2002
DEFINITION UI-E-CR0-acm-a-09-0-UI.rl UI-E-CR0 Homo sapiens cDNA clone
           UI-E-CF0-acm-a-09-0-UI 5', mRNA sequence.
ACCESSION
            BM689172
            BM689172.1 GI:19002430
VERSION
KEYWORDS
           EST.
SOURCE
           human.
  ORGANISM Homo sapiens
           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Euteleostomi;
           Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
REFERENCE
           1 (bases 1 to 357)
 AUTHORS
           Bonaldo, M. F., Lennon, G. and Soares, M. B.
 TITLE
           Normalization and subtraction: two approaches to facilitate
gene
           discovery
           Genome Res. 6 (9), 791-806 (1996)
  JOURNAL
           97044477
 MEDLINE
COMMENT
           Contact: Soares, MB
           Program for Rat Gene Discovery and Marping
           University of Iowa
            451 Eckstein Medical Research Building Iowa City, IA 52242,
USA
           Tel: 319 335 8250
           Fax: 319 335 9565
           Email: msoares@blue.weeq.uiowa.edu
           Tissue Procurement: Dr. Gregg Hageman
             DNA Library preparation: Dr. M. Bento Soares, Univeristy
of Iowa
```

```
cDNA Library Arrayed by: Dr. M. Bento Scares, Univeristy of
Towa
             DNA Sequencing by: Dr. M. Bento Soares, Univeristy of Iowa
             Clone Distribution: Researchers may obtain clones from
Research
            Genetics (www.resgen.com).
            Seq primer: M13 Reverse.
FEATURES
                     Location/Qualifiers
                     1. .357
     source
                      /organism="Homo sapiens"
                      /db xref="taxon:9606"
                      /clone="UI-E-CRO-acm-a-09-0-UI"
                      /clone lib="UI-E-CRO"
                      /tissue type="eye anterior segment"
                      /dev stage="adult"
                      /lab host="DH10B (Life Technologies) (T1 phage
resistant)"
                      /note="Organ: eye; Vector: pT7T3-Pac (Pharmacia)
with a
                      modified polylinker; Site 1: EcoR I; Site 2: Not I;
                      UI-E-CRO is a cDNA library containing the following
                      tissue(s): eye anterior segment. The library was
                      constructed according to Bonaldo, Lennon and
Scares,
                      Genome Research, 6:791-806, 1996. First strand cDNA
                      synthesis was primed with an oligo-dT primer
containing a
                      Not I site. Double stranded cDNA was ligated to an
ECOR T
                      adaptor, digested with Not I, and cloned
directionally
                      into pT7T3-Pac vector. The oligonucleotide used to
prime
                      the synthesis of first-strand cDNA contains a
library tag
                      sequence that is located between the Not I site and
the
                      (dT)18 tail. The sequence tag for this library is
                      AATGCCGCAT. This library was created for the
program, Gene
                      Discovery in the Visual System, supported by
National Eye
                      Institute (NEI)."
BASE COUNT
                           88 c 100 g
                 92 a
                                              77 t
ORIGIN
 Query Match 50.0; Score 20; DB 14; Length 357; Best Local Similarity 100.0; Pred. No. 0.72; Matches 20; Conservative 0; Mismatches 0; Indels
Gars
       21 GTATGGTGTGTGGATGCGAG 40
Qγ
          213 GTATGGTGTGTGGATGCGAG 232
FIGURE E
```

SEQ ID NO: 6

## **RESULT 5**

US-08-488-013-1/c

; Sequence 1, Application US/08488013

; Patent No. 5707806

: GENERAL INFORMATION:

; APPLICANT: Shuber, Anthony P.

; TITLE OF INVENTION: Direct Sequence Identification of

; Patent No. 5707806

; TITLE OF INVENTION: Mutations by Cleavage-and Ligation-Associated

; TITLE OF INVENTION: Mutation-Specific Sequencing

; NUMBER OF SEQUENCES: 1

; CORRESPONDENCE ADDRESS:

ADDRESSEE: Darby & Darby P.C.

STREET: 805 Third Avenue, 27th Floor

; CITY: New York City

STATE: New York

COUNTRY: USA

ZIP: 10022

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentln Release #1.0, Version #1.30

; CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/488,013

; FILING DATE: 07-JUN-1995

CLASSIFICATION: 435

; ATTORNEY/AGENT INFORMATION:

NAME: Ludwig, S. Peter

REGISTRATION NUMBER: 25,351

REFERENCE/DOCKET NUMBER: 0372/0B127

; TELECOMMUNICATION INFORMATION:

TELEPHONE: (212) 527-7770

TELEFAX: (212) 753-6237

TELEX: 236687

: INFORMATION FOR SEQ ID NO: 1:

; SEQUENCE CHARACTERISTICS:

```
LENGTH: 30 base pairs
   TYPE: nucleic acid
   STRANDEDNESS: single
   TOPOLOGY: linear
  MOLECULE TYPE: other nucleic acid
   DESCRIPTION: /desc = "synthetic oligonucleotide"
; HYPOTHETICAL: NO
; ANTI-SENSE: NO
US-08-488-013-1
                 50.0%; Score 20; DB 1; Length 30;
 Query Match
 Best Local Similarity 100.0%; Pred. No. 0.0096;
 Matches 20; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy
     1 GCGGTCCCAAAAGGGTCAGT 20
     Db
    30 GCGGTCCCAAAAGGGTCAGT 11
FIGURE F
SEQ ID NO: 7
RESULT 9
US-09-056-285A-1
; Sequence 1, Application US/09056285A
; Patent No. 6403307
    GENERAL INFOFMATION:
         AFPLICANT: Stone, Edwin M.
                     Sheffield, Val C.
                     Alward, Wallace L.M.
                     Fingert, John
         TITLE OF INVENTION: GLAUCOMA THERAPEUTICS AND DIAGNOSTICS
         NUMBER OF SEQUENCES: 43
         CORRESPONDENCE ADDRESS:
              ADDRESSEE: FOLEY, HCAG & ELIOT LLP
              STREET: One Post Office Square
              CITY: Boston
              STATE: MA
              COUNTRY: USA
              ZIF: 02109-2170
         COMPUTER READABLE FORM:
              MEDIUM TYPE: Floppy disk
              COMPUTER: IBM PC compatible
              OPERATING SYSTEM: PC-DOS/MS-DOS
              SOFTWARE: PatentIn Release #1.0, Version #1.30
         CURRENT APPLICATION DATA:
              APPLICATION NUMBER: US/09/056,285A
```

FILING DATE: 07-Apr-1998

```
ATTOFNEY/AGENT INFOFMATION:
;
             NAME: Arnold, Eeth E.
             REGISTRATION NUMBER: 35,430
             REFERENCE/LOCKET NUMBER: UIA-010.28
        TELECOMMUNICATION INFORMATION:
             TELEPHONE: 617-832-1000
;
             TELEFAM: 617-832-7000
;
    INFORMATION FOR SEQ ID NO: 1:
        SEQUENCE CHAFACTERISTICS:
;
             LENGTH: 2800 base pairs
             TYPE: nucleic acid
             STRANDEDNESS: single
             TOPOLOGY: linear
        MOLECULE TYPE: DNA (genomic)
        SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-056-285A-1
                         53.74; Score 22; DB 4; Length 2800;
 Query Match
  Best Local Similarity 100.0-; Pred. No. 0.00035;
 Matches 22; Conservative 0; Mismatches
                                               0; Indels
                                                                0;
Gaps
       Ú;
       20 TGCGAATAGAGCCATAAACTCA 41
         872 TGCGAATAGAGCCATAAACTCA 893
Db
FIGURE G
SEQ ID NO: 8
RESULT 7
US-09-277-016-16
; Sequence 16, Application US/09277016
; Patent No. 6143529
; GENERAL INFORMATION:
  APPLICANT: Lapidus, Stanley N
  APPLICANT: Shuber, Anthony P
  TITLE OF INVENTION: Methods for improving sensitivity and specificity
;
οf
  TITLE OF INVENTION: screening assays
  FILE FEFERENCE: EMT-030
  CUPRENT APPLICATION NUMBER: US/09/277,016
  CUFFENT FILING DATE: 1999-03-26
  EAFLIER AFFLICATION NUMBER: 08/876,857
;
  EAPLIEF FILING DATE: 1997-06-16
  EAFLIEF AFFLICATION NUMBEF: 08/700,583
  EAFLIER FILING DATE: 1996-08-14
  NUMBER OF SEQ ID NOS: 37
  SOFTWAFE: Patentin Ver. 2.0
; SEQ ID NO 16
   LENGTH: 37
;
   TYPE: DNA
   OFGANISM: Artificial Sequence
   FEATURE:
   OTHEF INFOFMATION: Description of Artificial Sequence: PCR-E-FOR (p53
   OTHER INFORMATION: Exon 5)
US-09-177-016-16
                         50.0-; Score 20; DB 3; Length 37;
  Query Match
  Best Local Similarity 100.0:, Pred. No. 0.0096;
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